Liver proteomics unravel the metabolic pathways related to Feed Efficiency in beef cattle

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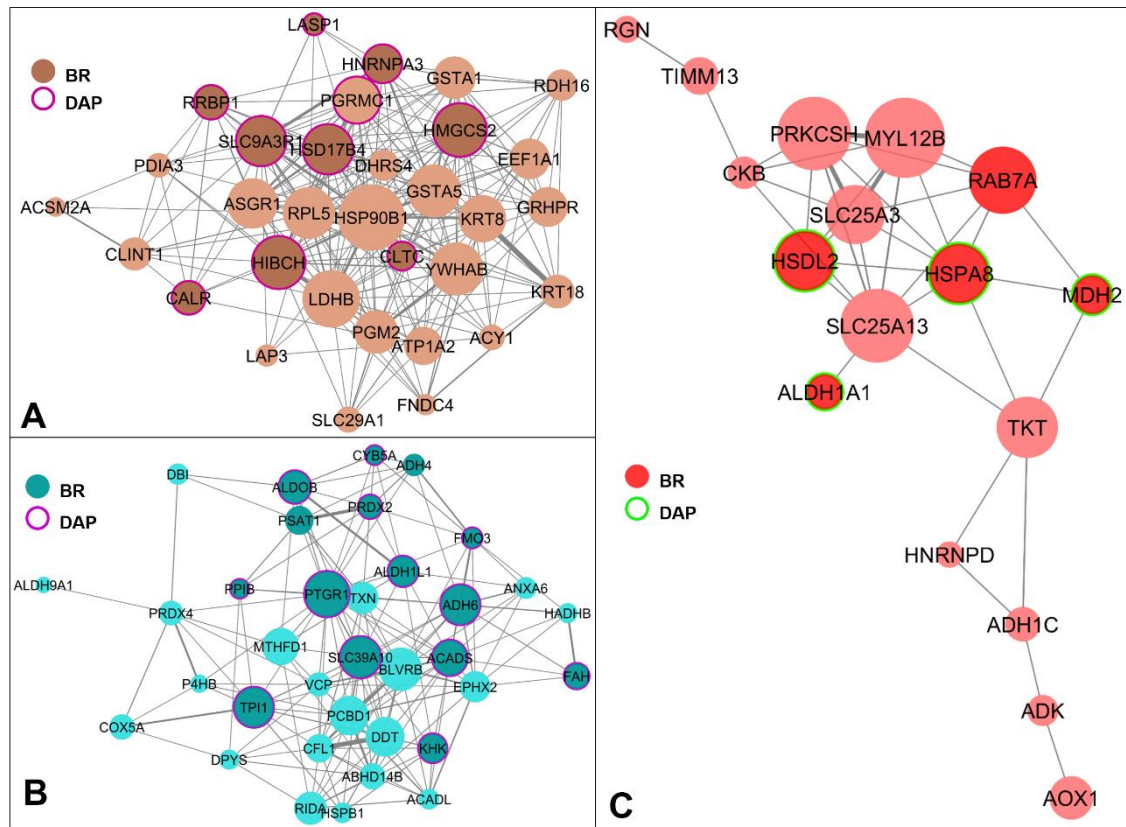
Figure S1 – Connectivity of hepatic proteins of Nellore cattle in co-expression networks related with feed efficiency. A: Connectivity in the brown module. B: Connectivity in the turquoise module. C: Connectivity in the red module. BR: biologically relevant proteins for feed efficiency, considering protein significance value > |0.60| (p-value < 0.05) for feed efficiency; DAP: differentially abundant proteins between Nellore cattle of high and low feed efficiency. Modules named by different colors represent co-expressed proteins network. Nodes represent the differentially abundant proteins that are identified with the coding gene symbol. Lines represent the connections between proteins.